

591508035Seqlist.txt
SEQUENCE LISTING

<110> KURODA, Masaharu
<120> Plant with Reduced Protein Content in Seed, Method of
Constructing the Same and Method of Using the Same

<130> 59150-8035

<140> US 10/539,992
<151> 2003-12-09

<150> PCT/JP2003/015753
<151> 2003-12-09

<150> JP 2002-369700
<151> 2002-12-20

<160> 119

<170> PatentIn version 3.3

<210> 1
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<213> Oryza sativa

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Leu Gln Pro His Leu Met Leu Gln Gln Gln Met Leu Ser Pro Cys Gly
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Glu Phe Val Arg Gln Gln Cys Ser Thr Val Ala Thr Pro Phe Phe Gln
50 55 60
Ser Pro Val Phe Gln Leu Arg Asn Cys Gln Val Met Gln Gln Gln Cys
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Cys Gln Gln Leu Arg Met Ile Ala Gln Gln Ser His Cys Gln Ala Ile

591508035Seqlist.txt

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Leu Gln Ser His Leu Leu Gln Gln Gln Val Leu Ser Pro Cys Ser 35
35 40 45
Glu Phe Val Arg Gln Gln His Ser Ile Val Ala Thr Pro Phe Trp Gln 50
50 55 60
Pro Ala Thr Phe Gln Leu Ile Asn Asn Gln Val Met Gln Gln Gln Cys 65
65 70 75 80
Cys Gln Gln Leu Arg Leu Val Ala Gln Gln Ser His Tyr Gln Ala Ile 85
85 90 95
Ser Ser Val Gln Ala Ile Val Gln Gln Leu Gln Leu Gln Gln Val Gly 100
100 105 110
Val Val Tyr Phe Asp Gln Thr Gln Ala Gln Ala Gln Leu Leu Ala 115
115 120 125
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<213> Oryza sativa

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Leu Gln Ser Pro Val Leu Leu Gln Gln Gln Val Leu Ser Pro Tyr Asn
35 40 45
Glu Phe Val Arg Gln Gln Tyr Gly Ile Ala Ala Ser Pro Phe Leu Gln
50 55 60
Ser Ala Ala Phe Gln Leu Arg Asn Asn Gln Val Trp Gln His Gln Ala
65 70 75 80
Gly Gly Gln Gln Ser Arg Tyr Gln Asp Ile Asn Ile Val Gln Ala Ile
85 90 95
Ala Tyr Glu Leu Gln Leu Gln Phe Gly Asp Leu Tyr Phe Asp Arg
100 105 110
Asn Gln Ala Gln Ala Leu Leu Ala Phe Asn Val Pro Ser Arg
115 120 125
Tyr Gly Ile Tyr Pro Arg Tyr Tyr Gly Ala Pro Ser Thr Ile Thr Thr
130 135 140
Leu Gly Gly Val Leu
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caagtattat acagaaaaat agaagatct agtgtcccg agcaatgaag atcatttcg 180

591508035Seqlist.txt

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ctgcattca	actgagaaac	aaccaagtct	ggcaacagct	cgcgtggtg	gccaacaat	420
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Leu Gln Ser Pro Val Leu Leu Gln Gln Gln Val Leu Ser Pro Tyr Asn
35 40 45
Glu Phe Val Arg Gln Gln Tyr Gly Ile Ala Ala Ser Pro Phe Leu Gln
50 55 60
Ser Ala Ala Phe Gln Leu Arg Asn Asn Gln Val Trp Gln Gln Leu Ala
65 70 75 80
Leu Val Ala Gln Gln Ser His Tyr Gln Asp Ile Asn Ile Val Gln Ala
85 90 95
Ile Ala Gln Gln Leu Gln Leu Gln Phe Gly Asp Leu Tyr Phe Asp
100 105 110
Arg Asn Leu Ala Gln Ala Gln Leu Ala Phe Asn Val Pro Ser Arg Tyr
115 120 125
Gly Ile Tyr Pro Arg Tyr Tyr Gly Ala Pro Ser Thr Ile Thr Thr Leu
130 135 140
Gly Gly Val Leu
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attacagcga aagcataaca actagaatcc taccacaatg aagatcattt tcttctttgc 180
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Leu Gln Gln Gln Met Leu Ser Pro Cys Gly Glu Phe Val Arg Gln Gln
35 40 45
Cys Ser Thr Val Ala Thr Pro Phe Phe Gln Ser Pro Val Phe Gln Leu
50 55 60
Arg Asn Cys Gln Val Met Gln Gln Gln Cys Cys Gln Gln Leu Arg Met
65 70 75 80
Ile Ala Gln Gln Ser His Cys Gln Ala Ile Ser Ser Val Gln Ala Ile
85 90 95
Val Gln Gln Leu Gln Leu Gln Gln Phe Ser Gly Val Tyr Phe Asp Gln
100 105 110
Ala Gln Ala Gln Ala Gln Ala Met Leu Gly Leu Asn Leu Pro Ser Ile
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Cys Gly Ile Tyr Pro Ser Tyr Asn Thr Val Pro Glu Ile Pro Thr Val
130 135 140
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591508035Seqlist.txt

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Val	Val	Tyr	Phe	Asp	Gln	Thr	Gln	Ala	Gln	Ala	Gln	Ala	Leu	Leu	Ala
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Leu	Asn	Leu	Pro	Ser	Ile	Cys	Gly	Ile	Tyr	Pro	Asn	Tyr	Tyr	Ile	Ala
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<212> DNA

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Leu	Gln	Ser	His	Leu	Leu	Leu	Gln	Gln	Gln	Val	Leu	Ser	Pro	Cys	Ser
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				50		55			60						
Pro	Ala	Thr	Phe	Gln	Leu	Ile	Asn	Asn	Gln	Val	Met	Gln	Gln	Gln	Cys
				65		70			75		80				
Cys	Gln	Gln	Leu	Arg	Leu	Val	Ala	Gln	Gln	Ser	His	Tyr	Gln	Ala	Ile
				85		90			95						
Ser	Ile	Val	Gln	Ala	Ile	Val	Gln	Gln	Leu	Gln	Leu	Gln	Gln	Phe	Ser
				100		105			110						
Gly	Val	Tyr	Phe	Asp	Gln	Thr	Gln	Ala	Gln	Ala	Gln	Thr	Leu	Leu	Thr
				115		120			125						
Phe	Asn	Leu	Pro	Ser	Ile	Cys	Gly	Ile	Tyr	Pro	Asn	Tyr	Tyr	Ser	Ala

591508035Seqlist.txt

130	135	140
Pro Arg Ser Ile Ala Thr Val Gly Gly Val Trp Tyr		
145	150	155

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<213> Oryza sativa

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Val Gln Ser Pro Leu Leu Gln Gln Gln Val Leu Ser Pro Tyr Asn
35 40 45
Glu Phe Val Arg Gln Gln Tyr Ser Ile Ala Ala Ser Thr Phe Leu Gln
50 55 60
Ser Ala Ala Phe Gln Leu Arg Asn Asn Gln Val Leu Gln Gln Leu Arg
65 70 75 80
Leu Val Ala Gln Gln Ser His Tyr Gln Asp Ile Asn Val Val Gln Ala
85 90 95
Ile Ala His Gln Leu His Leu Gln Gln Phe Gly Asn Leu Tyr Ile Asp
100 105 110
Arg Asn Leu Ala Gln Ala Gln Ala Leu Leu Ala Phe Asn Leu Pro Ser
115 120 125
Thr Tyr Gly Ile Tyr Pro Trp Ser Tyr Ser Ala Pro Asp Ser Ile Thr
130 135 140
Thr Leu Gly Gly Val Leu Tyr
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591508035Seqlist.txt

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<223> 13kD prolamine

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Leu	Gln	Ser	Pro	Val	Leu	Leu	Gln	Gln	Gln	Val	Leu	Ser	Pro	Tyr	Asn
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Glu	Phe	Val	Arg	Gln	Gln	Tyr	Gly	Ile	Ala	Ala	Ser	Pro	Phe	Leu	Gln
						50		55			60				
Ser	Ala	Ala	Phe	Gln	Leu	Arg	Asn	Asn	Gln	Val	Trp	Gln	Gln	Leu	Ala
						65		70		75		80			
Leu	Val	Ala	Gln	Gln	Ser	His	Tyr	Gln	Asp	Ile	Asn	Ile	Val	Gln	Ala
						85		90		95					
Ile	Ala	Gln	Gln	Leu	Gln	Gln	Phe	Gly	Asp	Leu	Tyr	Phe	Asp		
						100		105		110					
Arg	Asn	Leu	Ala	Gln	Ala	Gln	Ala	Leu	Leu	Ala	Phe	Asn	Val	Pro	Ser
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cccatataat	gagttcgtaa	gcagcagttat	ggcataccggc	aacccttct	tgcaatcagc	240
tgcgttcaa	ctgagaaaca	accaagtctg	gcaacaggtc	gcgcgtgg	cgcaacaatc	300
tcactatcg	gacattaaaca	ttgttcaggc	catagcgcag	cagtcacaac	tccagcagtt	360
tggtgcattc	tactttgatc	ggaatctggc	tcaagctaa	gctctgttgg	cttttaacgt	420
gccacccaaa	tatggtatct	acccttagta	ctatggtca	cccagttacca	ttaccaccct	480
tggcgtgtc	ttgtatgtt	tttaacagta	taatggtgg	aagttaaaaaa	taagctcaga	540
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591508035Seqlist.txt

<210> 20
<211> 148
<212> PRT
<213> Oryza sativa

<220>
<223> 13kD prolamine

<400> 20
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1 5 10 15
Ala Ser Arg Ser Leu Ile Leu Gly Gln Ser Tyr Arg Gln Tyr Gln Leu
20 25 30
Gln Ser Pro Val Leu Leu Gln Gln Gln Val Leu Ser Pro Tyr Asn Glu
35 40 45
Phe Val Ser Ser Ser Met Ala Tyr Gly Asn Pro Phe Leu Gln Ser Ala
50 55 60
Ala Phe Gln Leu Arg Asn Asn Gln Val Trp Gln Gln Leu Ala Leu Val
65 70 75 80
Ala Gln Gln Ser His Tyr Gln Asp Ile Asn Ile Val Gln Ala Ile Ala
85 90 95
Gln Gln Leu Gln Leu Gln Gln Phe Gly Asp Leu Tyr Phe Asp Arg Asn
100 105 110
Leu Ala Gln Ala Leu Leu Ala Phe Asn Val Pro Pro Lys Tyr
115 120 125
Gly Ile Tyr Pro Arg Tyr Tyr Gly Ala Pro Ser Thr Ile Thr Thr Leu
130 135 140
Gly Gly Val Leu
145

<210> 21
<211> 769
<212> DNA
<213> Oryza sativa

<220>
<223> 13kD prolamine

<220>
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<222> (11)..(11)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (109)..(109)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (207)..(207)
<223> n is a, c, g, or t

<400> 21
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gccgcgcccc atccccggtgc gcgacccatc gttcacacag ttcaagcatt atacagaaaa 180
atagaaagat ctatgttccgc gcagccatgc aagatcattt tcgtctttgc tctccttgct 240
attgctgcatt gcaggccctt gcccggatgg atgttttttag gtcaaagttt taggcaatat 300
cagctgcagt cgcctgtctt gctacagcaa caggtgcattt gcccataaa tgagttcgta 360
aggcagcagt atggcatagc ggcaagcccc ttcttgcattt cagctgcattt tcaactgaga 420
aataaccaag tctggcaaca tcaggtgtgtt ggccaacaat ctcgcataca ggacattaaac 480
attgttcagg ccatagcgtt cggatcacaat ctccagcaat ttgggtatctt ctactttgat 540

591508035Seqlist.txt

cggaatcagg	ctcaagctca	agctctattg	gcttttaacg	tgccatctag	atatggtac	600
tacccttagt	actatggtgc	accaggattc	attaccaccc	ttggcggtgt	cttgtaatgt	660
gttttaacag	tatagtggtt	cggaagttaa	aaataagctc	agatatcatc	atatgtgaca	720
tgtgaaactt	tgggtgatat	aaatagaaat	aaagttgcct	ttcatatattt		769

<210> 22
<211> 149
<212> PRT
<213> Oryza sativa

<220>
<223> 13kD prolamine

<400> 22
Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Ala Ala Cys Arg
1 5 10 15
Pro Leu Pro Ser Leu Met Phe Leu Gly Gln Ser Tyr Arg Gln Tyr Gln
20 25 30
Leu Gln Ser Pro Val Leu Leu Gln Gln Gln Val Leu Ser Pro Tyr Asn
35 40 45
Glu Phe Val Arg Gln Gln Tyr Gly Ile Ala Ala Ser Pro Phe Leu Gln
50 55 60
Ser Ala Ala Phe Gln Leu Arg Asn Asn Gln Val Trp Gln His Gln Ala
65 70 75 80
Gly Gly Gln Gln Ser Arg Tyr Gln Asp Ile Asn Ile Val Gln Ala Ile
85 90 95
Ala Tyr Glu Leu Gln Leu Gln Gln Phe Gly Asp Leu Tyr Phe Asp Arg
100 105 110
Asn Gln Ala Gln Ala Leu Leu Ala Phe Asn Val Pro Ser Arg
115 120 125
Tyr Gly Ile Tyr Pro Arg Tyr Tyr Gly Ala Pro Ser Thr Ile Thr Thr
130 135 140
Leu Gly Gly Val Leu
145

<210> 23
<211> 609
<212> DNA
<213> Oryza sativa

<220>
<223> 13kD prolamine

<400> 23
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aatatttaggc aatatcggt gcagtcgcct ctcctgctac agcaacaggt gcttagccta 180
tataatgagt tcgttaaggca gcagtatagc attgcggcaa gccccttctt gcaatcagct 240
gtgttcaac tgagaaacaa ccaagtcttg caacagctca ggctgggtggc gcaacaatct 300
caactaccagg acattaacgt tgtccaggcc atagcgcagc agctacacct ccagcagttt 360
ggcgatctct acattgacgg gaatctggct caagcgcac gactgttggc ttttaacttg 420
ccatctacat atggtatcta cccttaggtac tatagagcac cggtagtat taccaccctt 480
ggcggtgtct tgtactgaat tttcacaata ttgttagttcg gaagtgaaaaa tataaggcctc 540
aggtatcatc gtatgtgaca tgtgaaactt aaggtgatat aaatagaaat aaaattatct 600
ttcatatattt 609

<210> 24
<211> 150
<212> PRT
<213> Oryza sativa

<220>
<223> 13kD prolamine

591508035Seqlist.txt

<400> 24
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 1 5 10 15
 Thr Ala Gln Phe Asp Val Leu Gly Gln Asn Ile Arg Gln Tyr Gln Val
 20 25 30
 Gln Ser Pro Leu Leu Leu Gln Gln Val Leu Ser Leu Tyr Asn Glu
 35 40 45
 Phe Val Arg Gln Gln Tyr Ser Ile Ala Ala Ser Pro Phe Leu Gln Ser
 50 55 60
 Ala Val Phe Gln Leu Arg Asn Asn Gln Val Leu Gln Gln Leu Arg Leu
 65 70 75 80
 Val Ala Gln Gln Ser His Tyr Gln Asp Ile Asn Val Val Gln Ala Ile
 85 90 95
 Ala Gln Gln Leu His Leu Gln Gln Phe Gly Asp Leu Tyr Ile Asp Arg
 100 105 110
 Asn Leu Ala Gln Ala Gln Arg Leu Leu Ala Phe Asn Leu Pro Ser Thr
 115 120 125
 Tyr Gly Ile Tyr Pro Arg Tyr Tyr Arg Ala Pro Gly Ser Ile Thr Thr
 130 135 140
 Leu Gly Gly Val Leu Tyr
 145 150

<210> 25
 <211> 596
 <212> DNA
 <213> Oryza sativa

<220>
 <223> 13kD prolamine

<400> 25
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 atatcagctg cagtcgcctg tcctgctaca gcaacatgtg cttagcccat ataatgagtt 180
 cgtaaaggcag cagttatggca tagccggcaag ccccttcctg caatcagctg cgtttcaact 240
 gagaacaaac caagtctggc aacagctcgc gctggtgccg caacaatctc actatcagga 300
 cattaaacatt gttcaggcga tagcgcagca gctacaactc cagcagtttgcgtatctct 360
 ctgttgcattggc aatctggc aagctcaagc tctgttgct tttaaacgtgc catctagata 420
 tggatctac ccttagtact atggtgccacc cagtaccatt accacccttgcgggtgtctt 480
 gtaatgagtt ttaacagtat agtggttcgg aagataaaaaa taagctcaga tatcatcata 540
 tgtgacatgt gaaactttgg gtgatataaa tagaaaaaaa gttgtcttc atattt 596

<210> 26
 <211> 149
 <212> PRT
 <213> Oryza sativa

<220>
 <223> 13kD prolamine

<400> 26
 Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Ala Ala Cys Arg
 1 5 10 15
 Pro Leu Gln Phe Asp Val Leu Gly Gln Ser Tyr Arg Gln Tyr Gln Leu
 20 25 30
 Gln Ser Pro Val Leu Leu Gln Gln His Val Leu Ser Pro Tyr Asn Glu
 35 40 45
 Phe Val Arg Gln Gln Tyr Gly Ile Ala Ala Ser Pro Phe Leu Gln Ser
 50 55 60
 Ala Ala Phe Gln Leu Arg Asn Asn Gln Val Trp Gln Gln Leu Ala Leu
 65 70 75 80
 Val Ala Gln Gln Ser His Tyr Gln Asp Ile Asn Ile Val Gln Ala Ile

591508035Seqlist.txt

Ala Gln Gln Leu Gln Leu Gln Gln	85 Phe Gly Asp Leu Tyr Phe Asp Arg	90 95
100	105	110
Asn Leu Ala Gln Ala Gln Ala	Leu Leu Ala Phe Asn Val Pro Ser Arg	
115	120	125
Tyr Gly Ile Tyr Pro Arg Tyr	Gly Ala Pro Ser Thr Ile Thr Thr	
130	135	140
Leu Gly Gly Val Leu		
145		

<210> 27
<211> 285
<212> DNA
<213> Oryza sativa

<220>
<223> 13kD prolamine

<400> 27
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tttgataaac aaccaagtca tgcagcagca gttttgccaa cagtcaggc tggtagcaca 120
acatttcac taccaggcca tttagtattgt tcaagcgatt gtgcaacagc tacaactgca 180
gcatttttagt ggtgtctact ttgatcagac tcaagctcaa gcccaaactt ttttgacctt 240
caactttccc atccatatgtt ggtatctacc ttaacttact attgt 285

<210> 28
<211> 94
<212> PRT
<213> Oryza sativa

<220>
<223> 13kD prolamine

<400> 28
Phe Val Arg Gln Gln Tyr Ser Ile Val Ala Thr Pro Phe Trp Gln Pro
1 5 10 15
Ala Thr Phe His Leu Ile Asn Asn Gln Val Met Gln Gln Gln Phe Cys
20 25 30
Gln Gln Leu Arg Leu Val Ala Gln His Ser His Tyr Gln Ala Ile Ser
35 40 45
Ile Val Gln Ala Ile Val Gln Gln Leu Gln Leu Gln His Phe Ser Gly
50 55 60
Val Tyr Phe Asp Gln Thr Gln Ala Gln Ala Gln Thr Phe Leu Thr Phe
65 70 75 80
Asn Phe Pro Ser Ile Cys Gly Ile Tyr Leu Asn Leu Leu Leu
85 90

<210> 29
<211> 1836
<212> DNA
<213> Oryza sativa

<220>
<223> 13kD prolamine

<400> 29
tccacatggg acggggccaa ggtgaggaaa gcaagctgca caaaggattt aagttcttgt 60
aaacttgaaa ctcatttga gtgttatcc tagctaatat gatcccttca tcctagaata 120
taacaatcta gaatttagatg tgctatctaa acacatttga gtaggtatg tgtcatctaa 180
tcttagatat aatctaaaac ggaaggtgaa acggaggagg tacctacata gtaatggcat 240
gcctatgttg cttatattga cccgtgcagc tgagtatatg tgatggagac aaaagtact 300
ttcattatgg caccaaagga gattgttgg ggtgcctaat agaacatcg a tccaaatgac 360
acgacacact tagattctaa taggacatcc aagcaaaaca acacttagat cctaatagga 420

591508035Seqlist.txt

catccaagca	aaactaacac	tctagagcaa	ccgataagga	attgaaaaag	tttgtccatc	480
attctgaca	agaggttagt	tacaaaaaaa	atatttagt	gagctctgc	tcactacgca	540
tcacagaagt	ataacctaga	tataattaat	tcagtataga	agcaaaaatt	cagcagaac	600
aatagggtt	aaaactagaa	agaaggattt	atgatgtcc	tcagttatt	cagtgc当地	660
agatagttt	ctgtaaaacaa	aatggataat	aaacctgtat	tttcaacaaa	actagaggaa	720
ctctgtaaat	tgtccagggt	catccctaga	agttggttc	tccttacggg	aggaggagtt	780
atatgtatg	gacacaaaag	ttactttcat	gatgaaacca	aagggtattt	gttggggcac	840
ctaacagaac	atctatctaa	atgacatgac	tcacttagat	cctaatacgaa	catccaagca	900
aaactaacac	tctaaagcaa	ccgatgagga	attgaaagaa	aatatatgcc	atcgcatcta	960
taaatagaca	agcccaatga	aaaccctcct	catcggttac	acagttcaag	cattatacag	1020
aaaagaagat	ctagtgtccc	gcagcaatga	agatcatttt	ccgtcttgc	tctccttgct	1080
attgctgcat	gcaacaccctc	tgcgttagtt	atgttttagg	tcaaagttat	aggcaatatc	1140
agctacagtc	gcctctccta	caacaacaac	aggtgcttag	cccatataat	gacttcgtaa	1200
ggcagcgata	tgccatagcg	gcaagccct	tcttgcatac	agctgcgtt	aaactgagaa	1260
ataaccaagt	ctggcaacag	ctcgggctgg	tggcgc当地	atctcactat	caggacatta	1320
acattgttca	ggccatagcg	cagcagctat	aactccagca	gtttgggtat	ctctactttg	1380
atcggaatcc	ggctcaagct	caagctctgt	tggctttaa	cgtgccc当地	agatatggta	1440
tctaccctag	gtactatagt	acacccagta	ccattaccac	ccttggcggt	gtcttgtaat	1500
gagtttaac	agtatagttg	ttcgaagtt	aaaaataagc	tcatatatta	tcatatgtga	1560
catgtgaaat	ttgggggtgaa	ataaatcgaa	ataaagtgt	ctttcatatt	taaataccat	1620
gcctctataa	ggatataatcc	tagtacattg	tcgtaactaa	ttaccatcat	cggtactcta	1680
caattttact	gtgttcttac	attcgatccg	aagctactt	gttttaaga	tataaatggaa	1740
gcgtataaag	gatgtccgtc	ctttcattcc	aataagaaca	atgtaacatc	ctgaaaatgt	1800
gtcatttct	aatcctgcat	catccgact	ctttagt			1836

<210> 30

<211> 101

<212> PRT

<213> Oryza sativa

<220>

<223> 13kD prolamine

<400> 30

Met	Lys	Ile	Ile	Phe	Arg	Leu	Cys	Ser	Pro	Cys	Tyr	Cys	Cys	Met	Gln
1				5				10						15	
His	Leu	Cys	Val	Val	Asp	Val	Leu	Gly	Gln	Ser	Tyr	Arg	Gln	Tyr	Gln
				20				25						30	
Leu	Gln	Ser	Pro	Leu	Leu	Gln	Gln	Gln	Gln	Val	Leu	Ser	Pro	Tyr	Asn
				35				40						45	
Asp	Phe	Val	Arg	Gln	Arg	Tyr	Gly	Ile	Ala	Ala	Ser	Pro	Phe	Leu	Gln
				50				55						60	
Ser	Ala	Ala	Phe	Lys	Leu	Arg	Asn	Asn	Gln	Val	Trp	Gln	Gln	Leu	Gly
				65				70						75	
Leu	Val	Ala	Gln	Gln	Ser	His	Tyr	Gln	Asp	Ile	Asn	Ile	Val	Gln	Ala
				85				90						95	
Ile	Ala	Gln	Gln	Leu											
				100											

<210> 31

<211> 622

<212> DNA

<213> rice

<220>

<223> 16kD prolamine

<400> 31

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ctatggcaa	tgcccgacgc	agccgttat	gcaaccgatc	atgaaccgg	gcaatgagtt	180
cgtgaggcaa	cagtgccggcc	cgtgaggcc	accttggaaag	cgtcagcga	ggctacaact	240
gagcagctgc	caggtgtatgc	ggcagcaatg	ctgtcagcag	atgaggttgc	tggcgcaaca	300

591508035Seqlist.txt

atatcattgc	caggctattt	gcaccatgg	gtcgtctatc	atgcagcaag	tgcagttga	360
tgctggctt	gttggcgagc	cccaagctca	ggcccaggcc	caggtggctc	tcaatttgcc	420
ctccatgtgt	ggagtctacc	ctaggtactg	cagcactcca	tgcaaagttg	ctactgtca	480
ttgcggttct	tggtagtgtg	taccatcata	tatataatgt	tggataaata	aagtgtaca	540
catcatcg	tgtgtcatgt	aataaaattt	ggaatagtct	ttggctgttc	gtatgaataa	600
atgaaaatta	taacaaaaaa	aa				622

<210> 32
<211> 149
<212> PRT
<213> rice

<220>
<223> 16kD prolamine

<400> 32
Met Lys Ile Phe Val Ile Leu Ser Leu Leu Ala Leu Ala Ala Ser Ser
1 5 10 15
Ala Ser Ala Gln Phe Asp Ala Cys Thr Tyr Gly Gln Cys Gln Gln Gln
20 25 30
Pro Phe Met Gln Pro Ile Met Asn Pro Cys Asn Glu Phe Val Arg Gln
35 40 45
Gln Cys Ser Pro Met Ser Leu Pro Trp Lys Gln Ser Arg Arg Leu Gln
50 55 60
Leu Ser Ser Cys Gln Val Met Arg Gln Gln Cys Cys Gln Gln Met Arg
65 70 75 80
Leu Met Ala Gln Gln Tyr His Cys Gln Ala Ile Cys Thr Met Val Gln
85 90 95
Ser Ile Met Gln Gln Val Gln Phe Asp Ala Gly Phe Val Gly Glu Pro
100 105 110
Gln Ala Gln Ala Gln Ala Gln Val Ala Leu Asn Leu Pro Ser Met Cys
115 120 125
Gly Val Tyr Pro Arg Tyr Cys Ser Thr Pro Cys Lys Val Ala Thr Gly
130 135 140
His Cys Gly Ser Trp
145

<210> 33
<211> 562
<212> DNA
<213> Oryza sativa

<220>
<223> 10kD prolamine

<400> 33
cgtctacacc atctggaatc ttgtttaaca ctagtattgt agaatcagca atggcagcat 60
acaccagcaa gatcttgcc ctgtttgcct taattgctct ttctgcaagt gccactactg 120
caatcaccac tatgcagtat ttcccaccaa cattagccat gggcaccatg gatccgtgta 180
ggcagtagat gatgcaaacg ttggcatgg gtagctccac agccatgttc atgtcgcagc 240
caatggcgct cctgcagcag caatgttgc a tgcagctaca aggcatgtatg cctcagtgcc 300
actgtggcac cagttgccag atgatgcaga gcatgcaaca agttatttgt gctggactcg 360
ggcagcagca gatgatgaag atggcgatgc agatgcccata catgtgcaac atggcccttg 420
tcaacttcca actctcttcc ttgtgttgc tttgatcaaa cgttggttac atgtactcta 480
gtaataaggt gttgcatact atcgatgtca aacactagaa ataagaacca ttgaataaaa 540
tatcaatcat tttcagactt gc 562

<210> 34
<211> 134
<212> PRT
<213> Oryza sativa

<220>

591508035Seqlist.txt

<223> 10kD prolamine

<400> 34

Met Ala Ala Tyr Thr Ser Lys Ile Phe Ala Leu Phe Ala Leu Ile Ala
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Leu Ser Ala Ser Ala Thr Thr Ala Ile Thr Thr Met Gln Tyr Phe Pro
20 25 30
Pro Thr Leu Ala Met Gly Thr Met Asp Pro Cys Arg Gln Tyr Met Met
35 40 45
Gln Thr Leu Gly Met Gly Ser Ser Thr Ala Met Phe Met Ser Gln Pro
50 55 60
Met Ala Leu Leu Gln Gln Cys Cys Met Gln Leu Gln Gly Met Met
65 70 75 80
Pro Gln Cys His Cys Gly Thr Ser Cys Gln Met Met Gln Ser Met Gln
85 90 95
Gln Val Ile Cys Ala Gly Leu Gly Gln Gln Met Met Lys Met Ala
100 105 110
Met Gln Met Pro Tyr Met Cys Asn Met Ala Pro Val Asn Phe Gln Leu
115 120 125
Ser Ser Cys Gly Cys Cys
130

<210> 35

<211> 332

<212> DNA

<213> Oryza rufipogon

<220>

<223> 10kD prolamine

<400> 35

aattgtcttt tctgcaagtgc ccaactactgc aatcaccact atgcagtatt tcccaccaac 60
attagccatg ggcaccatgg atccgtgttag gcagtagatg atgcaaacgt tgggcattggg 120
tagctccaca gccatgttca tgtcgacgccc aatggcgctc ctgcagcagc aatgttgcat 180
gcagctacaa ggcattatgtc ctcagtgcacat ctgtggcacc agttgccaga tgatgcagag 240
catgcaccaa gttattttgtc ctggactcgg gcagcagcag atgatgaaga tggcgatgca 300
gatgccatac atgtgcaaca tggccctgt ca 332

<210> 36

<211> 110

<212> PRT

<213> Oryza rufipogon

<220>

<223> 10kD prolamine

<400> 36

Ile Ala Leu Ser Ala Ser Ala Thr Thr Ala Ile Thr Thr Met Gln Tyr
1 5 10 15
Phe Pro Pro Thr Leu Ala Met Gly Thr Met Asp Pro Cys Arg Gln Tyr
20 25 30
Met Met Gln Thr Leu Gly Met Gly Ser Ser Thr Ala Met Phe Met Ser
35 40 45
Gln Pro Met Ala Leu Leu Gln Gln Gln Cys Cys Met Gln Leu Gln Gly
50 55 60
Met Met Pro Gln Cys His Cys Gly Thr Ser Cys Gln Met Met Gln Ser
65 70 75 80
Met Gln Gln Val Ile Cys Ala Gly Leu Gly Gln Gln Gln Met Met Lys
85 90 95
Met Ala Met Gln Met Pro Tyr Met Cys Asn Met Ala Pro Val
100 105 110

<210> 37

591508035Seqlist.txt

<211> 349
<212> DNA
<213> Oryza longistaminata

<220>
<223> 10kD prolamine

<220>
<221> misc_feature
<222> (18)..(19)
<223> n is a, c, g, or t

<400> 37
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agtatttccc accaacattna gccatggca ccatggatcc gtgttaggcag tacatgatgc 120
aaacgttggg catggtagc tccacaacca tggtcatgtc gcagccaatg gcgctcctgc 180
agcagcaatg ttgcattgcag ctacaaggca tgatgcctca gtgccactgt ggcaccagt 240
gccagatgt gcagagcatg caacaagttg tttgtgttgg actcgggcag cagcagatga 300
tgcataatgtt gcaatgcag atgcataca tgtgcaacat ggccctgtt 349

<210> 38
<211> 116
<212> PRT
<213> Oryza longistaminata

<220>
<223> 10kD prolamine

<220>
<221> misc_feature
<222> (6)..(6)
<223> Xaa can be any naturally occurring amino acid

<400> 38
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1 5 10 15
Thr Thr Met Gln Tyr Phe Pro Pro Thr Leu Ala Met Gly Thr Met Asp
20 25 30
Pro Cys Arg Gln Tyr Met Met Gln Thr Leu Gly Met Gly Ser Ser Thr
35 40 45
Thr Met Phe Met Ser Gln Pro Met Ala Leu Leu Gln Gln Gln Cys Cys
50 55 60
Met Gln Leu Gln Gly Met Met Pro Gln Cys His Cys Gly Thr Ser Cys
65 70 75 80
Gln Met Met Gln Ser Met Gln Gln Val Val Cys Ala Gly Leu Gly Gln
85 90 95
Gln Gln Met Met Lys Met Ala Met Gln Met Pro Tyr Met Cys Asn
100 105 110
Met Ala Pro Val
115

<210> 39
<211> 343
<212> DNA
<213> Oryza rufipogon

<220>
<223> 10kD prolamine

<400> 39
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ttcccacca cattagccat gggcaccatg gatccgttgc ggcaggatcat gatgcaaacg 120
ttggccatgg gtagctccac agccatgttc atgtgcacgc caatggcgct cctgcacgc 180

591508035Seqlist.txt

caatgttgca	tgcagctaca	aggcatgatg	cctcagtgcc	actgtggcac	cagttgccag	240
atgatgcaga	gcatgcaaca	agttatttgt	gctggactcg	ggcagcagca	gatgatgaag	300
atggcgcatgc	agatgccata	catgtgcaac	atggccctg	tca	343	

<210> 40
<211> 113
<212> PRT
<213> Oryza rufipogon

<220>
<223> 10kD prolamine

<400> 40
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1 5 10 15
Thr Met Gln Tyr Phe Pro Pro Thr Leu Ala Met Gly Thr Met Asp Pro
20 25 30
Cys Arg Gln Tyr Met Met Gln Thr Leu Gly Met Gly Ser Ser Thr Ala
35 40 45
Met Phe Met Ser Gln Pro Met Ala Leu Leu Gln Gln Cys Cys Met
50 55 60
Gln Leu Gln Gly Met Met Pro Gln Cys His Cys Gly Thr Ser Cys Gln
65 70 75 80
Met Met Gln Ser Met Gln Gln Val Ile Cys Ala Gly Leu Gly Gln Gln
85 90 95
Gln Met Met Lys Met Ala Met Gln Met Pro Tyr Met Cys Asn Met Ala
100 105 110
Pro

<210> 41
<211> 339
<212> DNA
<213> Oryza rufipogon

<220>
<223> 10kD prolamine

<400> 41
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ccaccaacat tagccatggg caccatggat ccgtgttaggc agtacatgat gcaaacgttg 120
ggcatggta gctccacagc catgttcatg tcgcagccaa tggcgctcct gcagcagcaa 180
tggcatgc agctacaagg catgatgcct cagtgccact gtggcaccag ttgcccagatg 240
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<210> 42
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<212> PRT
<213> Oryza rufipogon

<220>
<223> 10kD prolamine

<400> 42
Phe Ala Leu Ile Ala Leu Ser Ala Ser Ala Thr Thr Ala Ile Thr Thr
1 5 10 15
Met Gln Tyr Phe Pro Pro Thr Leu Ala Met Gly Thr Met Asp Pro Cys
20 25 30
Arg Gln Tyr Met Met Gln Thr Leu Gly Met Gly Ser Ser Thr Ala Met
35 40 45
Phe Met Ser Gln Pro Met Ala Leu Leu Gln Gln Cys Cys Met Gln
50 55 60
Leu Gln Gly Met Met Pro Gln Cys His Cys Gly Thr Ser Cys Gln Met

591508035Seqlist.txt

65	70	75	80
Met Gln Ser Met Gln Gln Val Ile Cys Ala Gly Leu Gly Gln Gln			
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Met Met Lys Met Ala Met Gln Met Pro Tyr Met Cys Asn Met Ala Pro			
	100	105	110
Val			

<210> 43
<211> 343
<212> DNA
<213> *Oryza rufipogon*

<220>
<223> 10kD prolamine

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<220>
<221> misc_feature
<222> (19)..(19)
<223> n is a, c, g, or t
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<400> 43							
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atttccacc	aacatttagcc	atgggcacca	tgatccgtg	taggcagtac	atgatgcaaa		120
cgttggcat	ggtagctcc	acagccatgt	tcatgtcgca	gccaatggcg	ctcctgcagc		180
agcaatgttgc	catgcagcta	caaggcatga	tgcctcagtg	ccactgtggc	accagttgcc		240
agatgatgca	gagcatgcaa	caagttttt	gtgctggact	cgggcagcag	cagatgatga		300
agatggcgat	gcagatgcca	tacatgtgca	acatggcccc	tgt			343

<210> 44
<211> 114
<212> PRT
<213> Oryza rufipogon

<220>
<223> 10kD prolamine

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<220>
<221> misc_feature
<222> (6)..(6)
<223> xaa can be any naturally occurring amino acid
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<210> 45
<211> 533
<212> DNA
<213> *Oryza sativa*

591508035Seqlist.txt

<220>

<223> 10kD prolamine

<400> 45

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atgtcgacg	caatggcgt	cctgctgcag	caatgttgca	tgcagctaca	aggcatgatg	240
cctcaagtgc	actgtggcac	cagttgcccag	atgatgcaga	gcatgcaaca	agttatgtt	300
gctggactcg	ggcagcaga	gatgatgaag	atggcgatgc	agatgccata	catgtgcaac	360
atggcccctg	tcaacttcca	actctttcc	tgtggttgtt	gttgcataaa	cgttggttac	420
atgtactcta	gtaataaggt	gttgcatact	atcggtgtca	aacactagaa	ataagtacca	480
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<210> 46

<211> 134

<212> PRT

<213> Oryza sativa

<220>

<223> 10kD prolamine

<400> 46

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	20				25			30							
Pro	Thr	Leu	Ala	Met	Gly	Thr	Met	Asp	Pro	Cys	Arg	Gln	Tyr	Met	Met
	35			40			45								
Gln	Thr	Leu	Gly	Met	Gly	Ser	Ser	Thr	Ala	Met	Phe	Met	Ser	Gln	Pro
	50			55			60								
Met	Ala	Leu	Leu	Leu	Gln	Gln	Cys	Cys	Met	Gln	Leu	Gln	Gly	Met	Met
65				70			75			80					
Pro	Gln	Cys	His	Cys	Gly	Thr	Ser	Cys	Gln	Met	Met	Gln	Ser	Met	Gln
	85			90			95								
Gln	Val	Ile	Cys	Ala	Gly	Leu	Gly	Gln	Gln	Met	Met	Lys	Met	Ala	
	100			105			110								
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	115			120			125								
Ser	Ser	Cys	Gly	Cys	Cys										
	130														

<210> 47

<211> 940

<212> DNA

<213> rice

<220>

<223> 10kDa prolamine promoter

<400> 47

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taattataat	atcagttaaa	attgaaaata	atgcaacttc	atacttgcatt	gtgttcgtat	180
gtgcctgcct	aagaaatgt	tcttgtcata	atatgattac	atgaaatatg	tttacttcct	240
tcgtttctct	ttatgtgtaa	gataaagaac	tagatatgt	gaaagttagga	tagcaaagag	300
tatggccaaa	ctctaatttt	tgcttttattt	tttgggatgg	acccaaaatt	tgtttctccct	360
ttacttcttt	ccctttacaa	caatgttctt	tacttccaa	tcttattaac	aaaactccaa	420
atacatgcca	aactgcata	gtatgtatgc	tattaaggca	catttacaaa	gctccaagtt	480
tacctactca	atcattcaca	tatggcgatg	actcaaactc	ttaattgtt	tctgtgtaa	540
ctgtgacttg	tgtaaacat	tctacaagtc	ccatacgaat	tctgttcaca	aaagtttctt	600
tgtccagctc	ataatttaca	aaactgcaaa	atgccaagtc	aacttggcac	aaccttatca	660
tcatattttc	tttccacgca	ttaaggact	ggcagaatta	tcttgcgtt	gatattccaa	720

591508035Seqlist.txt

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tgtataaata	ggcccctagg	caaccattat	tccatcatcc	tcaacaatat	tgtctacacc	900
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<210> 48						
<211> 1351						
<212> DNA						
<213> rice						
<220>						
<223> GLUTELIN-B1 promoter						
<400> 48						
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agatatggat	tttctaagat	taattgattc	tctgtctaaa	aaaaaaaagt	attattgttat	120
taaatggaaa	aagaaaaagg	aaaagggga	tggcttctgc	tttttggct	gaaggcggcg	180
tgtggccagc	gtgctgcgtg	cggacagcga	gccaacacac	gacggagcag	ctacgacgaa	240
cgggggaccg	agtggaccgg	acgaggatgt	ggccttagac	gagtgcacaa	ggctagtgga	300
ctcggtcccc	gcbcgggtatc	ccgagtggtc	cactgtctgc	aaacacgatt	cacatagagc	360
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agatgtttt	ttattgttat	ataaacttagc	tttgtctgtc	tttgaactca	catcaattag	1320
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<210> 49						
<211> 852						
<212> DNA						
<213> Unknown						
<220>						
<223> Description of Unknown Organism:CaMV 35S gene promoter						
<400> 49						
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gagaagata	tatttctcaa	gatcagaagt	actattccag	tatggacgat	tcaaggcttg	240
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tcaaaggcca	ttggagtcaaa	gattcaaata	gaggaccaa	cagaactcgc	cgtaaagact	360
ggcgaacagt	tcatacagag	tctcttacga	ctcaatgaca	agaagaaaat	cttcgtcaac	420
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aaagatggac	ccccacccac	gaggagcatc	gtggaaaaag	aagacgttcc	aaccacgtct	720
tcaaagcaag	ttgattgtat	tgtatctcc	actgacgtaa	ggatgacgc	acaatcccac	780
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591508035Seqlist.txt

<210> 50
 <211> 1047
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic antisense sequence

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 tccactatcg gcgagacttt ctacacagcc atcggtccag acggccgcgc ttctgcgggc 120
 gatttgtta cgcccgacag tcccgctcc ggatcggacg attgcgtcgc atgcaccctg 180
 cgcggcaagct gcatcatcgaa aattggccgtc aaccaagctc tgatagagtt ggtcaagacc 240
 aatgcggagc atatacgccc ggagccgcgg cgatcctgcgaa agctccggat gcctccgctc 300
 gaagtagcgc gtctgctgtt ccatacaagc caaccacggc ctccagaaga agatgttggc 360
 gacctcgat tgggaatccc cgaacatcgcc tctgctccgg tcaatgaccg ctgttatgcg 420
 gccattgtcc gtcaggacat tggggagcc gaaatccgcg tgacacggat gccggacttc 480
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<210> 51
 <211> 67
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic antisense sequence

<400> 51
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 gtttcat 67

<210> 52
 <211> 15
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic antisense sequence

<400> 52
 atgaagatca ttttc 15

<210> 53
 <211> 15
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic control sequence

<400> 53
 ggatccccggg gtacc 15

<210> 54

591508035Seqlist.txt

<211> 1047
<212> DNA
<213> Unknown

<220>
<223> Description of Unknown Organism:hygromycin phosphotransferase gene

<400> 54

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cgttatgtt	atcggtcactt	tgcacggcc	gcgctccga	ttccggaaagt	gttgcacatt	240
ggggagttca	gcgagagcct	gacctattgc	atctccgc	gtgcacaggg	tgtcacattg	300
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caactggcaaa	ctgtgatgga	cgacaccgtc	agtgcgtccg	tcgcgcaggg	tctcgatgag	540
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atgttcgggg	atccccaata	cgaggtcgcc	aacatctct	tctggaggcc	gtggttggct	720
tgtatggagc	acgagacgcg	ctacttcgag	cggaggcattc	cgagacttgc	aggatcgccg	780
cggctccggg	cgtatatgt	ccgcatttgc	cttgaccaac	tctatcagag	cttggttgac	840
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gaatagagta	gatgccgacc	gtctagt				1047

<210> 55

<211> 265
<212> DNA
<213> Unknown

<220>
<223> Description of Unknown Organism:Nos terminator

<400> 55

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ggtgtcatct	atgttactag	atcgg				265

<210> 56

<211> 341
<212> PRT
<213> Artificial Sequence

<220>
<223> Modified HPT synthetic sequence

<400> 56

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								20		25			30		
Glu	Ser	Arg	Ala	Phe	Ser	Phe	Asp	Val	Gly	Gly	Arg	Gly	Tyr	Val	Leu
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Arg	Val	Asn	Ser	Cys	Ala	Asp	Gly	Phe	Tyr	Lys	Asp	Arg	Tyr	Val	Tyr
								50		55		60			
Arg	His	Phe	Ala	Ser	Ala	Ala	Leu	Pro	Ile	Pro	Glu	Val	Leu	Asp	Ile
								65		70		75		80	
Gly	Glu	Phe	Ser	Glu	Ser	Leu	Thr	Tyr	Cys	Ile	Ser	Arg	Arg	Ala	Gln

591508035Seqlist.txt

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Gln Thr Ser Gly Phe Gly Pro	Phe	Gly	Gly
130	135	140	140
Thr Thr Trp Arg Asp Phe Ile Cys	Ala	Ile	Asp
145	150	155	160
His Trp Gln Thr Val Met Asp Asp	Thr	Val	Ser
165	170	175	175
Ala Leu Asp Glu Leu Met Leu Trp	Ala	Glu	Asp
180	185	190	190
His Leu Val His Ala Asp Phe Gly	Ser	Asn	Asn
195	200	205	205
Gly Arg Ile Thr Ala Val Ile Asp Trp Ser	Glu	Ala	Met
210	215	220	Phe
Ser Gln Tyr Glu Val Ala Asn Ile Phe	Phe	Trp	Arg
225	230	235	Pro
Cys Met Glu Gln Gln Thr Arg Tyr Phe	Glu	Arg	Arg
245	250	255	His
Ala Gly Ser Pro Arg Leu Arg Ala Tyr	Pro	Glu	Leu
260	265	270	Asp
Gln Leu Tyr Gln Ser Leu Val Asp Gly	Asn	Phe	Asp
275	280	285	Ala
Ala Gln Gly Arg Cys Asp Ala Ile Val	Arg	Ser	Ala
290	295	300	Gly
Gly Arg Thr Gln Ile Ala Arg Arg Ser	Ala	Val	Trp
305	310	315	Thr
Cys Val Glu Val Leu Ala Asp Ser Gly	Asn	Arg	Arg
325	330	335	Pro
Pro Arg Ala Lys Glu			
	340		

<210> 57

<211> 2158

<212> DNA

<213> Artificial Sequence

<220>

<223> CAMV35S-Modified HPT-NOS synthetic sequence

<400> 57

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591508035Seqlist.txt

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tggaccgatg	gctgtgtaga	agtactcgcc	gatagtggaa	accgacgccc	cagcactcg	1860
ccgaggggcaa	aggaatagag	tagatgcca	ccgtctagtg	aatttcccg	atcggttcaa	1920
catttggcaa	taaagtttct	taagattgaa	tcctgttgc	ggtcttgcg	tgattatcat	1980
ataatttctg	ttgaattacg	ttaagcatgt	aataattaac	atgtaatgca	tgacgttatt	2040
tatgagatgg	gtttttatga	tttagagtccc	gcaattatac	atthaatacg	cgatagaaaaa	2100
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<210> 58
<211> 1757
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic promoter sequence

<400> 58

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aaatttaata	tatatatata	tatatatata	tatatatata	tatatatata	tatatatata	180
tatatatata	tatatatata	tatatatata	tatcacatca	gtctctgcac	aaagtgcac	240
ctgggctgct	tcaattataa	agccccattc	accacattt	ctagatagtc	gaaaagcacc	300
atcaatattt	agcttcaggt	attttgggt	gtgttgggt	tggattgatt	ctaataatata	360
ccaaatcaat	ataattcact	accaaaat	accatagcc	tcacaactt	attaatttt	420
gtagcttaag	atggtatata	taataaccaa	ttaacaactg	attctaattt	tactacggcc	480
cagtatgtac	caataaaaaa	caacgagtat	gtttcttcc	atcgtatcg	tacacagtac	540
aaaaaaacct	ggccagccctt	tcttgggctg	gggctctt	tcgaaaagg	acaaaacgta	600
cacggcagta	acgcccgttc	gctgcgtgtt	aacggccacc	aaccccgcc	tgagcaaaacg	660
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ttccttctt	tccttctcgc	gttttgcgtt	tggggacgga	ctccccaaac	cgccttccc	780
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cacggtgcgt	ctctcttattt	tgttaggtca	ctgttgggt	tgataggtac	actgtatgtt	1620
ttgtgttta	gttcgtgtat	ctaacaatatt	ggaataattt	gattgactga	tttctgtgt	1680
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<210> 59
<211> 926
<212> DNA
<213> Unknown

<220>

591508035Seqlist.txt

<223> Description of Unknown Organism:GUS gene partial fragment

<400> 59

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gccaactcct	accgtacctc	gcattaccct	tacgctgaag	agatgctcg	ctgggcagat	240
gaacatggca	tcgtgggtat	tgatgaaact	gctgctgtcg	gctttaacct	ctctttaggc	300
attgggttcg	aagcgggcaa	caagccgaaa	gaactgtaca	gcaagaggc	agtcaacggg	360
gaaactcagc	aagcgcaccc	acaggcgatt	aaagagctga	tagcgcgtga	caaaaaccac	420
ccaagcgtgg	tgatgtggag	tattgccaac	gaaccggata	cccgccgcga	agtgcacggg	480
aatatttcgc	cactggcgga	agcaacgcgt	aaactcgacc	cgacgcgtcc	gatcaccctgc	540
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gaatacggcg	tggatacgtt	agccgggctg	cactcaatgt	acaccgacat	gtggagtgaa	780
gagtatcagt	gtgcgtatgtat	ggatatgtat	caccgcgtct	ttgatgcgt	cagcgcgtc	840
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<210> 60

<211> 1198

<212> DNA

<213> Oryza sativa

<400> 60

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gttttggta	cagacatgga	gatgttgg	atgctatgaa	tagtcgatag	tttaaatttg	180
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atatggttat	actacgtgac	atataattcat	gagtgaggtt	cagagttttg	gttgcgttcc	360
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catgtttccc	cttctctgga	aaattttgtt	ggcaacagat	gccttctcct	tcttcagct	480
tctgcttctt	tagtcgtt	ggaggaagca	gcaaatagtt	gatgatatga	gaatccctca	540
catcggttag	gtgttaccaca	cgacttttt	attattattt	ttattattat	tattatttt	600
caaaaataaa	atagatca	ccctccacaa	caagtagagc	aagttggtga	gttattgtaa	660
agtttcacaa	agctaaattt	aaagttattt	cattaactt	tttcatattt	caaacaagag	720
tgtcaatgga	acaatgaaa	ccatatgaca	tactataatt	ttgtttttat	tattgaattt	780
atataattca	agagaataa	atccacatag	ccgtaaagtt	ctacatgtgg	tgcattacca	840
aaatataat	agtttacaaa	acatgacaag	cttagtttg	aaaatttgc	tccttattcac	900
attgacacat	aaagtggatg	atgagtcata	atattatttt	tcttgctacc	catcatgtat	960
atatgatagc	cacaaggta	ctttgtatgt	gataccaaag	aacatttttta	ggtgcaccta	1020
acagaatatc	caaataat	gactcactt	gatcataata	gagcatcaag	taaaactaac	1080
actctaaagc	aaccgtatgg	aaagcatcta	taaatagaca	agcacaatga	aaatccctat	1140
catccttcac	cacaattcaa	atattatgt	tgaagcatag	tagtagaattc	caacaaca	1198

<210> 61

<211> 163

<212> DNA

<213> rice

<220>

<223> 10kDa prolamine terminator

<400> 61

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tagaaataag	aaccattgaa	taaaatata	atcatattca	gacttgcaaa	tattgggtat	120
ttggatttct	gtccccatgtc	cctcttgaaa	gccatgctgt	aca		163

<210> 62

<211> 984

<212> DNA

591508035Seqlist.txt

<213> Oryza sativa

<220>

<223> GLUTELIN-A3 promoter

<400> 62

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tgtataagaa	tttttaggtt	gtgtgcata	taagtgttagc	ttcttata	tttagtgc	180
actatcttca	caagcacatg	ctatagtatt	gttccaagat	gaaagaataa	ttcatcctt	240
ctaccaactt	gcatgatatt	atatttgcata	atatcctatc	tcttggctt	taatgaaatg	300
tgctgctggg	ttataccttgc	ccatggatt	tgagagac	ttgtatagct	gaaaccaacg	360
tatatgcgag	catggacaaa	gagaacaaa	tgcaaggatt	tttttata	gttcatg	420
cctggatggg	ttaatatcg	gatcatcaaa	aaagatatgc	ataaaattaa	agtaataa	480
ttgctcataa	gaaaccaaaa	ccaaaagcac	atatgtccta	aacaaactgc	attttgc	540
tcatgttagc	atacaagaga	taatata	cgtggatt	acttattc	tttttg	600
tccaaaatgt	agttaggtct	actgatttgc	taaagtgtat	tgcttactgt	agaagt	660
tccaaaagc	aatcactaaa	gcaacacaca	acgtatagtc	cacccgtcac	gtattctt	720
gtggaaagata	acaagaaggc	tcactgaaa	ataaaagca	agaaaaggat	atcaa	780
ccatttgtt	tcccattgt	acttgtatgt	ctat	atccac	tgtgtac	840
acttctatct	agttagtac	ttcatatgt	gacattaaca	aacttatct	taacat	900
tcgatcacta	ctttacttca	ctataaaagg	accaacat	atcaccat	ctcacaa	960
cattgagttc	agtcccacaa	aaac				984

<210> 63

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Synthetic antisense sequence

<400> 63

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30

<210> 64

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 64

atgaagatca ttttcgtatt tgctctcctt gctattttgc catgc

45

<210> 65

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 65

caaagttata gacaatataca actacaatcg

30

<210> 66

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

591508035Seqlist.txt

<223> Synthetic antisense sequence	
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<210> 67	
<211> 45	
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<223> Synthetic antisense sequence	
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<211> 45	
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<210> 69	
<211> 15	
<212> DNA	
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<400> 69	15
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<210> 70	
<211> 30	
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<210> 71	
<211> 16	
<212> DNA	
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<223> Synthetic antisense sequence	
<400> 71	16
tgcagcagca gtgttg	
<210> 72	
<211> 23	
<212> DNA	

591508035Seqlist.txt

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 72

tgccggcgtttccaa cag

23

<210> 73

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 73

Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Val Ala Cys Asn

1 5 10 15

Ala Ser Ala Arg Phe Asp

20

<210> 74

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 74

Met Lys Ile Ile Phe

1 5

<210> 75

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 75

Met Lys Ile Ile Phe Val Phe Ala Leu Leu

1 5 10

<210> 76

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic antisense sequence

<400> 76

Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Val Ala

1 5 10

<210> 77

<211> 10

<212> PRT

<213> Artificial Sequence

591508035Seqlist.txt

<220>
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Gln Ser Tyr Arg Gln Tyr Gln Leu Gln Ser
1 5 10

<210> 78
<211> 5
<212> PRT
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<220>
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<400> 78
Glu Phe Val Arg Gln
1 5

<210> 79
<211> 15
<212> PRT
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<220>
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<400> 79
Glu Phe Val Arg Gln Gln His Ser Ile Val Ala Thr Pro Phe Trp
1 5 10 15

<210> 80
<211> 15
<212> PRT
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<220>
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<400> 80
Gln Gln Ser His Tyr Gln Ala Ile Ser Ser Val Gln Ala Ile Val
1 5 10 15

<210> 81
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
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<400> 81
Ala Gln Ala Gln Ala
1 5

<210> 82
<211> 10
<212> PRT
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<220>
<223> Synthetic antisense sequence

591508035Seqlist.txt

<400> 82
Tyr Phe Asp Gln Thr Gln Ala Gln Ala Gln
1 5 10

<210> 83
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
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<400> 83
Gln Gln Gln Cys Cys
1 5

<210> 84
<211> 7
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<220>
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<400> 84
Gln Gln Gln Cys Cys Gln Gln
1 5

<210> 85
<211> 9
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Glu Phe Val Arg Gln Gln Cys Ser Pro
1 5

<210> 86
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<220>
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Cys Gln Val Met Gln Gln Gln Cys Cys Gln Gln
1 5 10

<210> 87
<211> 6
<212> PRT
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<220>
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<400> 87
Gln Gln Cys Cys Gln Gln
1 5

591508035Seqlist.txt

<210> 88
<211> 6
<212> PRT
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<220>
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<400> 88
Glu Phe Val Arg Gln Gln
1 5

<210> 89
<211> 144
<212> PRT
<213> Oryza sativa

<220>
<223> RM4

<400> 89
Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Ala Ala Cys Ser
1 5 10 15
Ala Ser Ala Gln Phe Asp Val Leu Gly Gln Ser Tyr Arg Gln Tyr Gln
20 25 30
Leu Gln Ser Pro Val Leu Leu Gln Gln Gln Val Leu Ser Pro Tyr Asn
35 40 45
Glu Phe Val Arg Gln Gln Tyr Gly Ile Ala Ala Ser Pro Phe Leu Gln
50 55 60
Ser Ala Ala Phe Gln Leu Gln Leu Ala Leu Val Ala Gln Gln Ser
65 70 75 80
His Tyr Gln Asp Ile Asn Ile Val Gln Ala Ile Ala Gln Gln Leu Gln
85 90 95
Leu Gln Gln Phe Gly Asp Leu Tyr Phe Asp Arg Asn Leu Ala Gln Ala
100 105 110
Gln Ala Leu Ala Phe Asn Val Pro Ser Arg Tyr Gly Ile Tyr Pro
115 120 125
Arg Tyr Tyr Gly Ala Pro Ser Thr Ile Thr Thr Leu Gly Gly Val Leu
130 135 140

<210> 90
<211> 156
<212> PRT
<213> Oryza sativa

<220>
<223> RM5

<400> 90
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1 5 10 15
Ala Ser Ala Arg Phe Asp Ala Leu Ser Gln Ser Tyr Arg Gln Tyr Gln
20 25 30
Leu Gln Ser His Leu Leu Leu Gln Gln Gln Val Leu Ser Pro Cys Ser
35 40 45
Glu Phe Val Arg Gln Gln His Ser Ile Val Ala Thr Pro Phe Trp Gln
50 55 60
Pro Ala Thr Phe Gln Leu Ile Asn Asn Gln Val Met Gln Gln Gln Cys
65 70 75 80
Cys Gln Gln Leu Arg Leu Val Ala Gln Gln Ser His Tyr Gln Ala Ile
85 90 95
Ser Ser Val Gln Ala Ile Val Gln Gln Leu Gln Leu Gln Gln Val Gly

591508035Seqlist.txt

100	105	110
Val Val Tyr Phe Asp Gln Thr Gln	Ala Gln Ala Gln Ala	Leu Leu Ala
115	120	125
Leu Asn Leu Pro Ser Ile Cys Gly Ile Tyr Pro Asn	Tyr Tyr Ile Ala	
130	135	140
Pro Arg Ser Ile Pro Thr Val Gly Gly Val Trp	Tyr	
145	150	155

<210> 91
 <211> 158
 <212> PRT
 <213> Oryza sativa

<220>
 <223> RM7

<400> 91			
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20	25	30	
Leu Gln Ser His Leu Leu Leu Gln Gln Gln Val Leu Ser Pro Cys Ser			
35	40	45	
Glu Phe Val Arg Gln Gln Tyr Ser Ile Val Ala Thr Pro Phe Trp Gln			
50	55	60	
Pro Ala Thr Phe Gln Leu Ile Asn Asn Gln Val Met Gln Gln Gln Arg			
65	70	75	80
Met Cys Cys Gln Gln Leu Arg Leu Val Ala Gln Gln Ser His Tyr Gln			
85	90	95	
Ala Ile Ser Ile Val Gln Ala Ile Val Gln Gln Leu Gln Leu Gln Gln			
100	105	110	
Phe Ser Gly Val Tyr Phe Asp Gln Thr Gln Ala Gln Ala Gln Thr Leu			
115	120	125	
Leu Thr Phe Asn Leu Pro Ser Ile Cys Gly Ile Tyr Pro Asn Tyr Tyr			
130	135	140	
Ser Ala Pro Arg Ser Ile Ala Thr Val Gly Gly Val Trp Tyr			
145	150	155	

<210> 92
 <211> 134
 <212> PRT
 <213> Oryza sativa

<220>
 <223> RM10

<400> 92			
Met Ala Ala Tyr Thr Ser Lys Ile Phe Ala Leu Phe Ala Leu Ile Ala			
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Leu Ser Ala Ser Ala Thr Thr Ala Ile Thr Thr Met Gln Tyr Phe Pro			
20	25	30	
Pro Thr Leu Ala Met Gly Thr Met Asp Pro Cys Arg Gln Tyr Met Met			
35	40	45	
Gln Thr Leu Gly Met Gly Ser Ser Thr Ala Met Phe Met Ser Gln Pro			
50	55	60	
Met Ala Leu Leu Gln Gln Cys Cys Met Gln Leu Gln Gly Met Met			
65	70	75	80
Pro Gln Cys His Cys Gly Thr Ser Cys Gln Met Met Gln Ser Met Gln			
85	90	95	
Gln Val Ile Cys Ala Gly Leu Gly Gln Gln Met Met Lys Met Ala			
100	105	110	
Met Gln Met Pro Tyr Met Cys Asn Met Ala Pro Val Asn Phe Gln Leu			
115	120	125	

591508035Seqlist.txt

Ser Ser Cys Gly Cys Cys
130

<210> 93
<211> 149
<212> PRT
<213> Oryza sativa

<220>
<223> RM16

<400> 93
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Ala Ser Ala Gln Phe Asp Ala Cys Thr Tyr Gly Gln Cys Gln Gln Gln
20 25 30
Pro Phe Met Gln Pro Ile Met Asn Pro Cys Asn Glu Phe Val Arg Gln
35 40 45
Gln Cys Ser Pro Met Ser Leu Pro Trp Lys Gln Ser Arg Arg Leu Gln
50 55 60
Leu Ser Ser Cys Gln Val Met Arg Gln Gln Cys Cys Gln Gln Met Arg
65 70 75 80
Leu Met Ala Gln Gln Tyr His Cys Gln Ala Ile Cys Thr Met Val Gln
85 90 95
Ser Ile Met Gln Gln Val Gln Phe Asp Ala Gly Phe Val Gly Glu Pro
100 105 110
Gln Ala Gln Ala Gln Ala Gln Val Ala Leu Asn Leu Pro Ser Met Cys
115 120 125
Gly Val Tyr Pro Arg Tyr Cys Ser Thr Pro Cys Lys Val Ala Thr Gly
130 135 140
His Cys Gly Ser Trp
145

<210> 94
<211> 596
<212> DNA
<213> Oryza sativa

<220>
<223> RM4

<400> 94
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atatacgtcgt cagtcgcctg tcctgctaca gcaacaggtg cttagccat ataatgagtt 180
cgtaaggcag cagttatggca tagccgcaag ccccttctt caatcagctg cgtttcaact 240
gagaaacaac caagtctggc aacagctcgc gctggtgccg caacaatctc actatcagga 300
cattaaacatt gttcaggcca tagcgcagca gctacaactc cagcagttt gttgatctcta 360
ctttgatcggt aatctggctc aagctcaagc tctgttgct tttaacgtgc catctagata 420
tggtatctac cctaggtact atgggtgcacc cagtaccatt accaccctt gcggtgtctt 480
gtaatgagtt ttaacagtt agtggttcg aagttaaaaa taagctcaga tatcatatat 540
gtgacatgtg aaactttggg tggatataat agaaaaaaag ttgtctttca tattta 596

<210> 95
<211> 597
<212> DNA
<213> Oryza sativa

<220>
<223> RM5

<400> 95
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Page 33

591508035Seqlist.txt

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tcaaagtat	agacaatatac	aactacaatc	gcatctcctg	ctacagcaac	aagtgcctag	180
cccatgcagt	gagttcgtaa	ggcaacagca	tagcatatgt	gcaacccct	tctggcaacc	240
agctacgtt	caattgataa	acaaccaagt	catgcagcaa	cagtgttgc	aacagctcag	300
gctggtagcg	caacaatctc	actaccaggc	cattagtagc	gttcaggcga	ttgtgcagca	360
actacagctg	cagcagggtcg	gtgttgcata	ctttgatcag	actcaagctc	aagctcaagc	420
tttgcggcc	ttaaacttgc	catccatata	tggtatctat	cctaactact	acattgctcc	480
gaggagcatt	cccaccgtt	gtgggtctg	gtactgaatt	gtaatagtat	aatggtcaa	540
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<210> 96
<211> 609
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<220>
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caactacagt	cgcacatctct	actacagcaa	caagtgccta	gcccatgcag	tgagttcgta	180
aggcaacagt	atagcatagt	ggcaacccccc	ttctggcaac	cagctacgtt	tcaattgata	240
aacaaccaag	tcatgcagca	gcagttgtgc	caacagctca	ggctggtagc	acaacaatct	300
cactaccagg	ccatttagtat	tgttcaagcg	attgtgcac	agctacaact	gcagcaattt	360
agtgggtct	actttgatca	gactcaagct	caagccccaa	ctctgttgac	cttcaacttg	420
ccatccatata	gtggtatcta	ccctaactac	tatagtgtc	ccaggagcat	tgccactgtt	480
gggggtgtct	gttactgaat	tgttaacaata	taatagttcg	tatgttaaaa	ataaaagtcat	540
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cctaaaaaa						609

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<220>
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<400> 97						
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gactgtctgc	ctctctcagt	ttacttggat	gcattgacaa	catccttttt	tgctattact	120
cgtatttgct	ctatacgctt	tggcatatct	catgttggaa	tttgccttt	taatccaaaa	180
ttggatgtaa	ttgaaagaat	cctacgttgt	agttatttgg	attttggtgt	aaaaaaaat	240
agcctgtta	gaagaagcaa	aattggattt	agttaaaagg	atactagatg	gtgttatttg	300
gattttggtg	caaatacaat	taggaggtt	gttttattca	agttaaagtt	tgttttaaaa	360
aaattctcct	aaaaagatag	atactagatt	tgcatatatg	cattgaaaat	tacatctcg	420
cttggcggtt	atacttttag	tccctctaaa	ttgttcaatc	atttatgtat	aaaagggaaa	480
tcattttata	tcacaaagta	tttatgtatg	aaggggaaaa	atattctgca	tgggttgaa	540
caaaatacgt	ggattgggtt	agcctaaca	tacttggaaa	gggtatgtat	ttgatgttagt	600
gcccacatgg	tgtcgcttga	cattaaaacg	atatgcagtc	aggattgagg	aacattgctg	660
acaatttact	atcgctgtt	gtgttgacca	caataattca	gatgtaccat	cctatcttct	720
aactagaaag	atgcatggaa	gtttcttaca	ttatcccttgc	cacttggaaat	tttagtgaaa	780
tatcattaaa	acataaccac	ttactttgtt	gtgtatgtaa	ataaatgttt	tatcccttgg	840
aaagttgtat	attcatatata	tcttacagta	aattttattgt	ttttcttttc	atttatttct	900
aaattttaaac	cacccttttg	gtagcttaag	aaaaatttta	tttttgacag	tcctgttttc	960
tgttttca	ccccctccagg	aaaaccagct	actagtggat	cc		1002

<210> 98
<211> 37
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<213> Oryza sativa

591508035Seqlist.txt

<400> 98
Gln Val Met Gln Gln Gln Cys Cys Gln Gln Leu Arg Leu Val Ala Gln
1 5 10 15
Gln Ser His Tyr Gln Ala Ile Ser Ser Val Gln Ala Ile Val Gln Gln
20 25 30
Leu Gln Leu Gln Gln
35

<210> 99
<211> 34
<212> PRT
<213> Oryza sativa

<400> 99
Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Val Ala Cys Asn
1 5 10 15
Ala Ser Ala Arg Phe Asp Ala Leu Ser Gln Ser Tyr Arg Gln Tyr Gln
20 25 30
Leu Gln

<210> 100
<211> 26
<212> PRT
<213> Oryza sativa

<400> 100
Glu Phe Val Arg Gln Gln His Ser Ile Val Ala Thr Pro Phe Trp Gln
1 5 10 15
Pro Ala Thr Phe Gln Leu Ile Asn Asn Gln
20 25

<210> 101
<211> 31
<212> PRT
<213> Oryza sativa

<400> 101
Tyr Phe Asp Gln Thr Gln Ala Gln Ala Gln Ala Leu Leu Ala Leu Asn
1 5 10 15
Leu Gln Ser Ile Cys Gly Ile Tyr Pro Asn Tyr Tyr Ile Ala Pro
20 25 30

<210> 102
<211> 111
<212> DNA
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<220>
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<222> (1)..(111)

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<222> (6)..(6)
<223> n is a, c, g, or t

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591508035Seqlist.txt

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<220>
<221> misc_feature
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<223> n is a, c, g, or t

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<222> (103)..(105)
<223> n is a, c, g, or t

<400> 102
car gtn atg car car car tgy tgy car car nnn nnn nnn gtn gcn car 48
Gln Val Met Gln Gln Gln Cys Cys Gln Gln Xaa Xaa Xaa Val Ala Gln
1 5 10 15

car nnn cay tay car gcn atg nnn nnn gtn car gcn atg gtn car car 96
Gln Xaa His Tyr Gln Ala Met Xaa Xaa Val Gln Ala Met Val Gln Gln
20 25 30

nnn car nnn car car
Xaa Gln Xaa Gln Gln 35 111

<210> 103

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591508035Seqlist.txt

<211> 102
<212> DNA
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<220>
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<222> (1)..(102)

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<222> (85)..(87)
<223> n is a, c, g, or t

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<223> n is a, c, g, or t

591508035Seqlist.txt

<400> 103	48
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Met Lys Met Met Phe Val Phe Ala Xaa Xaa Ala Met Val Ala Cys Asn	
1 5 10 15	
gcn nnn gcn nnn tty gay gcn nnn nnn car nnn tay nnn car tay car	96
Ala Xaa Ala Xaa Phe Asp Ala Xaa Xaa Gln Xaa Tyr Xaa Gln Tyr Gln	
20 25 30	
nnn car	102
Xaa Gln	
<210> 104	
<211> 78	
<212> DNA	
<213> Oryza sativa	
<220>	
<221> CDS	
<222> (1)..(78)	
<220>	
<221> misc_feature	
<222> (9)..(12)	
<223> n is a, c, g, or t	
<220>	
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<222> (22)..(24)	
<223> n is a, c, g, or t	
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<222> (30)..(30)	
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<222> (39)..(39)	
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591508035Seqlist.txt

<222> (57)..(57)
<223> n is a, c, g, or t

<220>
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<222> (64)..(66)
<223> n is a, c, g, or t

<400> 104
gar tty gtn nnn car car cay nnn atg gtn gcn acn ccn tty tgg car 48
Glu Phe Val Xaa Gln Gln His Xaa Met Val Ala Thr Pro Phe Trp Gln
1 5 10 15

ccn gcn acn tty car nnn atg aay aay car 78
Pro Ala Thr Phe Gln Xaa Met Asn Asn Gln
20 25

<210> 105
<211> 93
<212> DNA
<213> Oryza sativa

<220>
<221> CDS
<222> (1)..(93)

<400> 105
tac ttt gat cag act caa gct caa gct caa gct ttg ctg gcc tta aac 48
Tyr Phe Asp Gln Thr Gln Ala Gln Ala Gln Ala Leu Leu Ala Leu Asn
1 5 10 15

ttg caa tcc ata tgt ggt atc tat cct aac tac tac att gct ccg 93
Leu Gln Ser Ile Cys Gly Ile Tyr Pro Asn Tyr Tyr Ile Ala Pro
20 25 30

<210> 106
<211> 1426
<212> DNA
<213> Oryza sativa

<400> 106
tttagcattaa tatttataga gtgataaagt catactacaa taaatcctta tatattaatt 60
gggggtcata ctagaagccc catattaatc ctacgagagg tagaaaacta gaaattatcg 120
cactagtcaa gttgcacttg gcctagagtc tcaattgttag tataaatgtat ataataattc 180
taaaaattaaa attagcaaat aacaagttca attaggttg aagccgtaat tctattttta 240
taatttaatc attcttaaat ttagaattac taaaaataaa ttattaatac agcgttgtac 300
ttgctgtaga gactcatata gttttacga cgatttaata atttcaaaaaa taaatacagg 360
aaattgctaa gtttgcatac taaaatataa tattgtcata atataataat tctaaaattc 420
aaattaataa ataccaagtt gatgtttat taaaatataa tagtatgtgc cgcacagctt 480
gatgcttagt cttagatctt taaccgtgc acgctgggtt aattagcgat ggtgcaggc 540
acgtacccaa atttcttcac tggatca actagagtag ttaaacgagg gcatgtgtatg 600
aaggcttagt atttgcattt ttccaaattat ccctgcataa gtcaggctac aatagcacct 660
ggactacatg cagggattac aaaataggtg gtaaccacat ttaccgcgtt aaccctatca 720
aattcaaata aattttaaaa gtaatttgat ttttttaata aattttgtat gttttctcaa 780
gctttatccc gtttaccgtg cttactgccc gaggcaatgg gaaaccctca ctagaagttg 840
cacctgttct tttctgtca ccatatcatg ttgaatcatg tgcgttgtt cctttcgaa 900
gaaccgattt actacatgac tcatcaattc cactttacatg atcaaaagggt ttgttatggg 960
ggcaatgcctt ttgtgaaattt aaattttat tttgcgtcac gtttatcta gttaaacact 1020
acctacatc cattacaaaa cctcattcca caaaacgtatg catctagata aaaaatatga 1080
catgtaaagt gatgtatgac tcatgtttat tatcaaaaaat cgataacaat caaatgat 1140
aggttagtaaa gtaccttga aatggcatgt ccaagttatgt gtagctccac ctagcacaat 1200
atcccaagtg atcatcataa aaggcataca aatacaagca gccgatgtatg cacacaagaa 1260
acaacacaaa ttgcacaaaa ccaaaagcaa ccgtgcctt gagcatagag atcatgtat 1320

591508035Seqlist.txt

tcccactata aatacaaatg caccatatca agatgctcct cacccttact gaaaaatcac 1380
 aaacatcaaa acgttataag agttctctag catccatcac atagcc 1426

<210> 107
 <211> 1008
 <212> DNA
 <213> Oryza sativa

<400> 107
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 agagaggagg aagaagagga ggggtgacaa tgatatgtgg gccatgtggc cccaccatt 120
 ttttaattca ttctttgtt gaaactgaca tgggggtccc atgagaattt ttattttcg 180
 gatcgaatttccacgtaa gctacgtcaa tgctacgtca gatgaagacc gagtcaaattt 240
 agccacgtaa ggcgcacgtc agccaaaacc accatccaa cccggaggg acctcatctg 300
 cactggtttt gatagtttag ggaccgttg tatctgggtt ttgcattgaa ggacgaaaat 360
 caaatttggtaa gacaagttaa gggaccttaa atgaacttat tccatttcaa aatattctgt 420
 gagccatata tccgtgggct tccaatccctc ctcaaattaa agggcctttt taaaatagat 480
 aattgccttc ttcaagtac ccataaaaagt acaaaaactac taccacaag caacatgcgc 540
 agttacacac attttctgca catttccacc acgtcacaat gagctaagag ttatccctag 600
 gacaatctca ttagtgtaga tacatccatt aatcttttat cagaggcaaa cgtaaagccg 660
 ctctttatgaa caaaaatagg tgacacaaaaa gtgttatctg ccacatcat aacttcagaa 720
 attaccacac accaagagaa aaataaaaaaa aaatctttt gcaagctcca aatcttgaa 780
 accttttca ctctttgcag cattgtactt ttgcctttt tccaaccgat ccatgtcacc 840
 ctcaagcttc tacttgatct acacgaagct caccgtgcac acaaccatgg ccacaaaaaac 900
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 gagaaaaaaa aacatataca ttcttagtga ttgtctgatt gatcatca 1008

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 <212> DNA
 <213> Oryza sativa

<220>
 <221> CDS
 <222> (1)..(72)

<400> 108
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 1 5 10 15
 ctt tct gca agt gcc act act gca
 Leu Ser Ala Ser Ala Thr Thr Ala 72
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<210> 109
 <211> 24
 <212> PRT
 <213> Oryza sativa

<400> 109
 Met Ala Ala Tyr Thr Ser Lys Ile Phe Ala Leu Phe Ala Leu Ile Ala
 1 5 10 15
 Leu Ser Ala Ser Ala Thr Thr Ala 20

<210> 110
 <211> 66
 <212> DNA
 <213> Oryza sativa

<220>
 <221> CDS

591508035Seqlist.txt

<222> (1)..(66)

<400> 110

atg aag atc att ttc gta ttt gct ctc ctt gct att gtt gca tgc aat
Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Val Ala Cys Asn
1 5 10 15

48

gct tct gca cggttt gat
Ala Ser Ala Arg Phe Asp
20

66

<210> 111

<211> 22

<212> PRT

<213> Oryza sativa

<400> 111

Met Lys Ile Ile Phe Val Phe Ala Leu Leu Ala Ile Val Ala Cys Asn
1 5 10 15
Ala Ser Ala Arg Phe Asp
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<210> 112

<211> 57

<212> DNA

<213> Oryza sativa

<220>

<221> CDS

<222> (1)..(57)

<400> 112

atg aag atc ttt gtc atc ctc tct ctc ctc gcc ctc gca gcg agc agc
Met Lys Ile Phe Val Ile Leu Ser Leu Leu Ala Leu Ala Ala Ser Ser
1 5 10 15

48

gcc tcg gca
Ala Ser Ala

57

<210> 113

<211> 19

<212> PRT

<213> Oryza sativa

<400> 113

Met Lys Ile Phe Val Ile Leu Ser Leu Leu Ala Leu Ala Ala Ser Ser
1 5 10 15
Ala Ser Ala

<210> 114

<211> 72

<212> DNA

<213> Oryza sativa

<220>

<221> CDS

<222> (1)..(72)

<400> 114

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Met Ala Ser Ser Val Phe Ser Arg Phe Ser Ile Tyr Phe Cys Val Leu
1 5 10 15

48

cta tta tgc cat ggt tct atg gcc
 Leu Leu Cys His Gly Ser Met Ala
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<210> 115
 <211> 24
 <212> PRT
 <213> Oryza sativa

<400> 115
 Met Ala Ser Ser Val Phe Ser Arg Phe Ser Ile Tyr Phe Cys Val Leu
 1 5 10 15
 Leu Leu Cys His Gly Ser Met Ala
 20

<210> 116
 <211> 66
 <212> DNA
 <213> Oryza sativa

<220>
 <221> CDS
 <222> (1)..(66)

<400> 116
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 1 5 10 15

gtg gcc atc tcc ggc gcg 66
 Val Ala Ile Ser Gly Ala
 20

<210> 117
 <211> 22
 <212> PRT
 <213> Oryza sativa

<400> 117
 Met Ala Ser Lys Val Val Phe Phe Ala Ala Ala Leu Met Ala Ala Met
 1 5 10 15
 Val Ala Ile Ser Gly Ala
 20

<210> 118
 <211> 8
 <212> PRT
 <213> Oryza sativa

<400> 118
 Ser Arg Ala Met Val Ser Leu Gly
 1 5

<210> 119
 <211> 102
 <212> DNA
 <213> recombinant construct

<400> 119
 atggcagcat acaccagcaa gatcttgcc ctgttgcct taattgctct ttctgcaagt 60
 gccactactg catctagagc aatggtgagc aaggcgagg ag 102